

01 JUL 2005

123 Sequence Listing.ST25
SEQUENCE LISTING

<110> Lee, Sang Yup
Lee, Seok Jae

<120> A PROTEIN CHIP FOR ANALYZING INTERACTION BETWEEN PROTEIN AND
SUBSTRATE PEPTIDE THEREOF

<130> 4240-123

<140> Not yet assigned
<141> 2005-07-01

<150> PCT/KR2003/002183
<151> 2003-10-18

<150> 10-2003-0000464
<151> 2003-01-04

<160> 12

<170> PatentIn version 3.2

<210> 1
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic Construct

<400> 1

Leu Arg Arg Ala Ser Leu Gly
1 5

<210> 2
<211> 33
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Construct

<400> 2
cggaattcat atggtgccca tccaaaaagt cca

33

<210> 3
<211> 48
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Construct

<400> 3
gcggatcctt agcccaggct cgcacgacgc aggcacccag ggctgagg

48

<210> 4
<211> 53
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Construct

<400> 4

123 Sequence Listing.ST25
gcggatcctt agcccaggct cgcgcggcgc agggggccca ggctcgacg acg 53

<210> 5
<211> 53
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Construct

<400> 5
gcggatcctt agcccaggct cgcgcggcgc agggggccca ggctcgacg acg 53

<210> 6
<211> 47
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Construct

<400> 6
catgccatgg gcatcaccat catcaccatg atattcaaaa aagagtg 47

<210> 7
<211> 50
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Construct

<400> 7
gctctagatt agcccaggct cgcacgacgc aggatggagg tacggcggtg 50

<210> 8
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic Construct

<400> 8

Glu Ala Ile Tyr Ala Ala Pro Phe Ala Lys Lys
1 5 10

<210> 9
<211> 33
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Construct

<400> 9
cggaattcat atggtgccca tccaaaaagt cca 33

<210> 10
<211> 67
<212> DNA
<213> Artificial Sequence

123 Sequence Listing.ST25

<220>

<223> Synthetic Construct

<400> 10

cgggatcctc attatTTTTT tttcgcaaac ggcgccgcat agatcgcttc gcaccaggg 60
ctgaggt 67

<210> 11

<211> 61

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Construct

<400> 11

cgggatcctt tttttttcgc aaacggcgcc gcatagatcg cttcgacccc agggctgagg 60
t 61

<210> 12

<211> 71

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Construct

<400> 12

cgggatcctc attatTTTTT tttcgcaaac ggcgccgcat agatcgcggg ttttttttc 60
gcaaacggcg c 71